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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/694,777A

DATE: 04/03/2002

TIME: 11:00:21

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3 <110> APPLICANT: PARDO-FERNANDEZ, LUIS ANGEL
4 STUHMER, WALTER
5 BECKH, SYNNOVE
6 BRUGGEMANN, ANDREA
7 FERNANDEZ-MIRANDA, DONATO DEL CAMINO
8 PEREZ, ARACELI SANCHEZ
9 WESELOH, RUDIGER
11 <120> TITLE OF INVENTION: NOVEL HUMAN K+ ION CHANNEL AND THERAPEUTIC APPLICATIONS
12 THEREOF
14 <130> FILE REFERENCE: MPG-8
16 <140> CURRENT APPLICATION NUMBER: 09/694,777A
17 <141> CURRENT FILING DATE: 2000-10-23
19 <150> PRIOR APPLICATION NUMBER: PCT/EP99/02695
20 <151> PRIOR FILING DATE: 1999-04-21
22 <150> PRIOR APPLICATION NUMBER: EP 98 10 7268.9
23 <151> PRIOR FILING DATE: 1998-04-21
25 <160> NUMBER OF SEQ ID NOS: 24
27 <170> SOFTWARE: PatentIn Ver. 2.1
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30 <211> LENGTH: 3002
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158 20 25 30
160 Gly Asn Ala Gln Ile Val Asp Trp Pro Ile Val Tyr Ser Asn Asp Gly

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169	Glu Lys Val Arg Gln Thr Phe Glu Asn Tyr Glu Met Asn Ser Phe Glu		
170	85	90	95
172	Ile Leu Met Tyr Lys Lys Asn Arg Thr Pro Val Trp Phe Phe Val Lys		
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175	Ile Ala Pro Ile Arg Asn Glu Gln Asp Lys Val Val Leu Phe Leu Cys		
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178	Thr Phe Ser Asp Ile Thr Ala Phe Lys Gln Pro Ile Glu Asp Asp Ser		
179	130	135	140
181	Cys Lys Gly Trp Gly Lys Phe Ala Arg Leu Thr Arg Ala Leu Thr Ser		
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184	Ser Arg Gly Val Leu Gln Gln Leu Ala Pro Ser Val Gln Lys Gly Glu		
185	165	170	175
187	Asn Val His Lys His Ser Arg Leu Ala Glu Val Leu Gln Leu Gly Ser		
188	180	185	190
190	Asp Ile Leu Pro Gln Tyr Lys Gln Glu Ala Pro Lys Thr Pro Pro His		
191	195	200	205
193	Ile Ile Leu His Tyr Cys Val Phe Lys Thr Thr Trp Asp Trp Ile Ile		
194	210	215	220
196	Leu Ile Leu Thr Phe Tyr Thr Ala Ile Leu Val Pro Tyr Asn Val Ser		
197	225	230	235
199	Phe Lys Thr Arg Gln Asn Asn Val Ala Trp Leu Val Val Asp Ser Ile		
200	245	250	255
202	Val Asp Val Ile Phe Leu Val Asp Ile Val Leu Asn Phe His Thr Thr		
203	260	265	270
205	Phe Val Gly Pro Ala Gly Glu Val Ile Ser Asp Pro Lys Leu Ile Arg		
206	275	280	285
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209	290	295	300
211	Pro Tyr Asp Val Ile Asn Ala Phe Glu Asn Val Asp Glu Gly Ile Ser		
212	305	310	315
214	Ser Leu Phe Ser Ser Leu Lys Val Val Arg Leu Leu Arg Leu Gly Arg		
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217	Val Ala Arg Lys Leu Asp His Tyr Ile Glu Tyr Gly Ala Ala Val Leu		
218	340	345	350
220	Val Leu Leu Val Cys Val Phe Gly Leu Ala Ala His Trp Met Ala Cys		
221	355	360	365
223	Ile Trp Tyr Ser Ile Gly Asp Tyr Glu Ile Phe Asp Glu Asp Thr Lys		
224	370	375	380
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 238 Glu Lys Ile Phe Ala Val Ala Ile Met Met Ile Gly Ser Leu Leu Tyr
 239 450 455 460
 241 Ala Thr Ile Phe Gly Asn Val Thr Thr Ile Phe Gln Gln Met Tyr Ala
 242 465 470 475 480
 244 Asn Thr Asn Arg Tyr His Glu Met Leu Asn Ser Val Arg Asp Phe Leu
 245 485 490 495
 247 Lys Leu Tyr Gln Val Pro Lys Gly Leu Ser Glu Arg Val Met Asp Tyr
 248 500 505 510
 250 Ile Val Ser Thr Trp Ser Met Ser Arg Gly Ile Asp Thr Glu Lys Val
 251 515 520 525
 253 Leu Gln Ile Cys Pro Lys Asp Met Arg Ala Asp Ile Cys Val His Leu
 254 530 535 540
 256 Asn Arg Lys Val Phe Lys Glu His Pro Ala Phe Arg Leu Ala Ser Asp
 257 545 550 555 560
 259 Gly Cys Leu Arg Ala Leu Ala Met Glu Phe Gln Thr Val His Cys Ala
 260 565 570 575
 262 Pro Gly Asp Leu Ile Tyr His Ala Gly Glu Ser Val Asp Ser Leu Cys
 263 580 585 590
 265 Phe Val Val Ser Gly Ser Leu Glu Val Ile Gln Asp Asp Glu Val Val
 266 595 600 605
 268 Ala Ile Leu Gly Lys Gly Asp Val Phe Gly Asp Val Phe Trp Lys Glu
 269 610 615 620
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 272 625 630 635 640
 274 Asp Leu His Val Ile Lys Arg Asp Ala Leu Gln Lys Val Leu Glu Phe
 275 645 650 655
 277 Tyr Thr Ala Phe Ser His Ser Phe Ser Arg Asn Leu Ile Leu Thr Tyr
 278 660 665 670
 280 Asn Leu Arg Lys Arg Ile Val Phe Arg Lys Ile Ser Asp Val Lys Arg
 281 675 680 685
 283 Glu Glu Glu Arg Met Lys Arg Lys Asn Glu Ala Pro Leu Ile Leu
 284 690 695 700
 286 Pro Pro Asp His Pro Val Arg Arg Leu Phe Gln Arg Phe Arg Gln Gln
 287 705 710 715 720
 289 Lys Glu Ala Arg Leu Ala Ala Glu Arg Gly Gly Arg Asp Leu Asp Asp
 290 725 730 735
 292 Leu Asp Val Glu Lys Gly Asn Val Leu Thr Glu His Ala Ser Ala Asn
 293 740 745 750
 295 His Ser Leu Val Lys Ala Ser Val Val Thr Val Arg Glu Ser Pro Ala
 296 755 760 765
 298 Thr Pro Val Ser Phe Gln Ala Ala Ser Thr Ser Gly Val Pro Asp His
 299 770 775 780
 301 Ala Lys Leu Gln Ala Pro Gly Ser Glu Cys Leu Gly Pro Lys Gly Gly
 302 785 790 795 800
 304 Gly Gly Asp Cys Ala Lys Arg Lys Ser Trp Ala Arg Phe Lys Asp Ala
 305 805 810 815
 307 Cys Gly Lys Ser Glu Asp Trp Asn Lys Val Ser Lys Ala Glu Ser Met

VERIFICATION SUMMARY

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